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P.Z.RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/672,665DATE: 02/06/2002
TIME: 07:23:50Input Set : A:\Cura-721.app
Output Set: N:\CRF3\02062002\I672665.raw

3 <110> APPLICANT: Prayaga, Sudhirdas K
 4 Vernet, Corine
 5 Shimkets, Richard A
 6 Burgess, Catherine E
 7 Spytek, Kimberly A
 9 <120> TITLE OF INVENTION: Novel Polynucleotides and Polypeptides Encoded Thereby
 11 <130> FILE REFERENCE: 15966-72, 76, 76A, 80, 112
 13 <140> CURRENT APPLICATION NUMBER: 09/672,665
 14 <141> CURRENT FILING DATE: 2000-09-28
 16 <150> PRIOR APPLICATION NUMBER: 60/156,745
 17 <151> PRIOR FILING DATE: 1999-09-30
 19 <150> PRIOR APPLICATION NUMBER: 60/158,942
 20 <151> PRIOR FILING DATE: 1999-10-06
 22 <150> PRIOR APPLICATION NUMBER: 60/159,248
 23 <151> PRIOR FILING DATE: 1999-10-13
 25 <150> PRIOR APPLICATION NUMBER: 60/169,344
 26 <151> PRIOR FILING DATE: 1999-12-06
 28 <150> PRIOR APPLICATION NUMBER: 60/215,048
 29 <151> PRIOR FILING DATE: 2000-06-29
 31 <160> NUMBER OF SEQ ID NOS: 32
 33 <170> SOFTWARE: PatentIn Ver. 2.0
 35 <210> SEQ ID NO: 1
 36 <211> LENGTH: 327
 37 <212> TYPE: DNA
 38 <213> ORGANISM: Homo sapiens
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 44 gaggaagagg aggagaaaaa agaagggtat ggtgaggaag aggtatggaga tgaagaggaa 240
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 58 20 25 30
 60 Ala Pro Ala Asn Gly Asn Ala Asn Glu Asn Gly Glu Gln Glu Ala
 61 35 40 45

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131 gacgcagccg tagacaccag ctccgaaatc accaacaagg acttaaagga gaagaaggaa 120
 132 gttgtggaaag aggcaaaaaa tgaaagagac gcccctgcta acggaaatgc taatgaggaa 180
 133 aatggggagc aggaggctga caatgaggtt gacgaagaag aggaagaagg tgggaggaa 240
 134 gaaggtgatg gtgaggaaga gatggagat gaagatgagg aagctgagtc agctacggc 300
 135 aagcgggcag ctgaagatga tgaggataac gatgtcgata ccaagaagca gaagaccgac 360
 136 gaggatgacc agacggcaaa aaaggaaaag ttaaactaaa aaaaaaaaaag gccggcgtga 420
 137 cctattcacc ctccacttcc cgtctcagaa tctaaacgtg gtcaccttcg agtagagagg 480
 138 cccgcccggc caccgtggc agtgccaccc gcagatgaca cgcgctctcc accacccaac 540
 139 ccaaaccatg agaatttgca acagggagg aaaaaagaac caaaacttcc aaggcctgct 600
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 153 20 25 30
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 156 35 40 45
 158 Asp Asn Glu Val Asp Glu Glu Glu Glu Gly Gly Glu Glu Glu Gly
 159 50 55 60
 161 Asp Gly Glu Glu Glu Asp Gly Asp Glu Glu Ala Glu Ser Ala
 162 65 70 75 80
 164 Thr Gly Lys Arg Ala Ala Glu Asp Asp Glu Asp Asn Asp Val Asp Thr
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 182 gaggaaaatg gggagccgga ggctgacaac gaggtatgtt aagaagagga agaagggtggg 180
 183 gaggagaaga gttatgtgtt ggaagaggat ggatgttgaatggggagc tgagtctgtt 240
 184 acgggcacggc gggcagctgtt agatgtatgtt gatgtatgtt tcgatataccca gaagcagaag 300
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 189 <212> TYPE: PRT
 190 <213> ORGANISM: Homo sapiens
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196 Leu Lys Glu Lys Lys Glu Val Val Glu Glu Ala Glu Asn Gly Arg Asp
197 20 25 30
199 Ala Pro Ala His Gly Asn Ala Asn Glu Glu Asn Gly Glu Pro Glu Ala
200 35 40 45
202 Asp Asn Glu Val Asp Glu Glu Glu Glu Gly Gly Glu Glu Glu Gly
203 50 55 60
205 Asp Gly Glu Glu Glu Asp Gly Asp Glu Asp Glu Gly Ala Glu Ser Ala
206 65 70 75 80
208 Thr Gly Lys Arg Ala Ala Glu Asp Asp Glu Asp Asp Asp Val Asp Thr
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226 ggtgtccgg accatgtcag acgcagccgt agacaccgc tccgaaatca ccaccaagga 180
227 cttaaagaag aaggaagctg tggaggaagc ggaaaatgga agagacaccc ctgctaatgg 240
228 gaaggctaat gaggaaaatg gggagcagga agctgacaat gaagttagatg aagaagagga 300
229 agaaggtggg gaggaagacg aggaggaaga agaaggcgat ggtgaggaag aggatggta 360
230 tgaagacgag gaagctgagt ccgctacggt caagcggca gctgaagatg atgagaatga 420
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244 20 25 30
246 Pro Ala Asn Gly Lys Ala Asn Glu Glu Asn Gly Glu Gln Glu Ala Asp
247 35 40 45
249 Asn Glu Val Asp Glu Glu Glu Glu Gly Glu Glu Asp Glu Glu
250 50 55 60
252 Glu Glu Glu Gly Asp Gly Glu Glu Asp Gly Asp Glu Asp Glu Glu
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 270 gaggaaaatg gggagccgga ggctgacaac gaggttagatg aagaagagga agaaggtggg 180
 271 gaggaagaag gtgatggtga ggaagaggat ggagatgaag atgagggagc tgagtcaagct 240
 272 acgggcaagc gggcagctga agatgtatgag gataacgatg tcgataccca gaagcagaag 300
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 282 1 5 10 15
 284 Leu Lys Glu Lys Lys Glu Val Val Glu Glu Ala Glu Asn Gly Arg Asp
 285 20 25 30
 287 Ala Pro Ala His Gly Asn Ala Asn Glu Glu Asn Gly Glu Pro Glu Ala
 288 35 40 45
 290 Asp Asn Glu Val Asp Glu Glu Glu Glu Gly Gly Glu Glu Glu Gly
 291 50 55 60
 293 Asp Gly Glu Glu Glu Asp Gly Asp Glu Asp Glu Gly Ala Glu Ser Ala
 294 65 70 75 80
 296 Thr Gly Lys Arg Ala Ala Glu Asp Asp Glu Asp Asn Asp Val Asp Thr
 297 85 90 95
 299 Gln Lys Gln Lys Thr Asp Glu Asp Asp Gln Thr Ala Lys Lys Glu Lys
 300 100 105 110
 302 Leu Asn
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 307 <211> LENGTH: 324
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 314 gaagaagatg gggacctgca gtgcctgtgt gtgaagacca cctcccaggt ccgtcccagg 180
 315 cacatcacca gcctggaggt gatcaaggcc ggacccact gccccactgc ccaactgtg 240
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 331 Ser Ile Ser Pro Ser Ala Glu Ala Glu Glu Asp Gly Asp Leu Gln Cys

VERIFICATION SUMMARY

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